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=> s wd-40

101 WD

24971 40

L1 14 WD-40

(WD(W)40)

# => d 1-14 ti

- L1 ANSWER 1 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI A combination of the F-box motif and kelch repeats defines a large Arabidopsis family of F-box proteins.
- L1 ANSWER 2 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI Proteomic analysis of arabidopsis seed germination and priming.
- L1 ANSWER 3 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2003) on STN
- TI Discrete domains mediate the light-responsive nuclear and cytoplasmic localization of Arabidopsis COP1.
- L1 ANSWER 4 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI Identification of partners of TIF34, a component of the yeast elF3 complex, required for cell proliferation and translation initiation.
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- (2003) on STN
- TI A conserved family of WD-40 proteins binds to the retinoblastoma protein in both plants and animals.
- L1 ANSWER 6 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2003) on STN
- TI Insulin-induced maturation of Xenopus oocytes is inhibited by microinjection of a Brassica napus cDNA clone with high similarity to a mammalian receptor for activated protein kinase C.
- L1 ANSWER 7 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI A wat1 mutant of fission yeast is defective in cell morphology.
- L1 ANSWER 8 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI Expression of an N-terminal fragment of COP1 confers a dominant-negative effect on light-regulated seedling development in Arabidopsis.
- L1 ANSWER 9 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI The Polycomb-group gene, extra sex combs, encodes a nuclear member of the WD-40 repeat family.
- L1 ANSWER 10 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI Molecular cloning and characterization of cDNA for a rice protein that contains seven repetitive segments of the Trp-Asp forty-amino-acid repeat (WD-40 repeat).
- L1 ANSWER 11 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI Non-clathrin-coat protein alpha is a conserved subunit of coatomer and in Saccharomyces cerevisiae is essential for growth.
- L1 ANSWER 12 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI Isolation of cDNAs encoding guanine nucleotide-binding protein beta-subunit homologues from maize (ZGB1) and Arabidopsis (AGB1).
- L1 ANSWER 13 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI PAS7 encodes a novel yeast member of the WD-40 protein family essential for import of 3-oxoacyl-CoA thiolase, a PTS2-containing protein, into peroxisomes.
- L1 ANSWER 14 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States

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- of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI Genetic and molecular analysis of an allelic series of cop1 mutants suggests functional roles for the multiple protein domains.

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- L1 ANSWER 1 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

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- In the sequences released by the Arabidopsis Genome Initiative (AGI), we AB have discovered a new large gene family (48 genes as of July 2000). A detailed computational and biochemical analysis of the predicted gene products reveals a novel family of plant F-box proteins, where the amino (N)-terminal F-box motif is followed by four kelch repeats and a characteristic carboxy-terminal domain. F-box proteins are an expanding family of eukaryotic proteins, which have been shown in some cases to be critical for the controlled degradation of cellular regulatory proteins via the ubiquitin pathway. The F-box motif of the At5g48990 gene product, a member of the family, was shown to be functionally active by its ability to mediate the in vitro interaction between At5g48990 and ASK1 proteins. F-box proteins specifically recruit the targets to be ubiquitinated, mainly through protein-protein interaction modules such as WD-40 domains or leucine-rich repeats (LRRs). The kelch repeats of the family described here form a potential protein-protein interaction domain, as molecular modelling of the kelch repeats according to the galactose oxidase crystal structure (the only solved structure containing kelch repeats) predicts a beta-propeller. The identification of this family of F-box proteins greatly expands the field of plant F-box proteins and suggests that controlled degradation of cellular proteins via the ubiquitin pathway could play a critical role in multiple plant cellular processes.

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- SO Plant molecular biology, July 2001. Vol. 46, No. 5. p. 603-614 Publisher: Dordrecht: Kluwer Academic Publishers. CODEN: PMBIDB; ISSN: 0167-4412

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- L1 ANSWER 3 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- AB The Arabidopsis CONSTITUTIVE PHOTOMORPHOGENIC1 (COP1) protein plays a critical role in the repression of photomorphogenesis during Arabidopsis seedling development. We investigated the control of COP1 partitioning between nucleus and cytoplasm, which has been implicated in the regulation of COP1 activity, by using fusion proteins between COP1 and beta-glucuronidase or the green fluorescent protein. Transient expression assays using onion epidermal cells and data from hypocotyl cells of stably transformed Arabidopsis demonstrated that COP1 carries a single, bipartite nuclear localization signal that functions independently of light. Nuclear exclusion was mediated by a novel and distinct signal, bordering the zinc-finger and coiled-coil motifs, that was able to redirect a

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heterologous nuclear protein to the cytoplasm. The cytoplasmic localization signal functioned in a light-independent manner. Light regulation of nuclear localization was reconstituted by combining the individual domains containing the nuclear localization signal and the cytoplasmic localization signal; the WD-40 repeat domain of COP1 was not required. However, phenotypic analysis of transgenic seedlings suggested that the constitutively nuclear-localized WD-40 repeat domain was able to mimic aspects of COP1 function, as indicated by exaggerated hypocotyl elongation under light conditions.

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- L1 ANSWER 5 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- AB In mammalian cells, the retinoblastoma (RB) protein regulates G1 progression and functions through its association with various cellular proteins. Two closely related mammalian RB binding proteins. RbAp48 and RbAp46, share sequence homology with the Msil protein of yeast. MSIl is a multicopy suppressor of a mutation in the IRA1 gene involved in the Ras-cAMP pathway that regulates cellular growth. Human RbAp48 is present in protein Complexes involved in histone acetylation and chromatin assembly. We report the cloning of cDNAs encoding four plant RbAp48-and Msi1-like proteins: one from tomato, LeMSI1, and three from Arabidopsis. Complementation Studies confirm that LeMSI1 can function as a multicopy suppressor of the yeast iral mutant phenotype. The LeMSIl protein localizes to the nucleus and binds to a 65-kD protein in wild-type as well as ripening inhibitor (rin) and Neverripe (Nr) tomoto fruit. LeMSI1 also binds to the human RB protein and the RB-like RRB1 protein from maize, indicating that this interaction is conserved between plants and animals.

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- L1 ANSWER 5 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- TI A conserved family of WD-40 proteins binds to the retinoblastoma protein in both plants and animals.

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- SO The Plant cell, Sept 1997. Vol. 9, No. 9. p. 1595-1606
  Publisher: [Rockville, MD : American Society of Plant Physiologists, c1989CODEN: PLCEEW; ISSN: 1040-4651

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